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Title:
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Listing first 45 summaries
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             Pred. No. 1s the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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2: /cgn2_6/ptodata/1.

3: /cgn2_6/ptodata/1.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
            TELEFAX: (619) 535-8949
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                             FILLNY ..... 435
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
NAME: Campbell, Cathryn B.
NAME: Campbell, Cathryn B.
                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
COMPOTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                  REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: CD40 Associated Proteins NUMBER OF SEQUENCES: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Reed, John C. APPLICANT: Sato, Takaaki
                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
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CITY: San Diego
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4370 La Jolla Village Drive,
543 amino acids
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                                                                                                                                                                                                                                           Sequence 36, Application US/09268544B Patent No. 6410710 GENERAL INFORMATION:
                                                                                                SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 36
LENGTH: 438
Best Local Similarity Matches 416; Conserv
                             Query Match
                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/268,544B CURRENT FILING DATE: 1999-03-11 NUMBER OF SEQ ID NOS: 43
                                                                                                                                                                                     APPLICANT: Lederman, Seth
APPLICANT: van Eyndhoven, Winfried
TITLE OF INVENTION: TRAF-3 Deletion Isoforms And Uses Thereof
FILE REFERENCE: 0575-58732
                                                                       ORGANISM: Human
                                                                                       TYPE: PRT
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                71.1%;
73.2%;
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Pred. No. 3.9e-231;
7; Mismatches 16;
    Score 2129.5;
Pred. No. 5.7e
7; Mismatches
                   7e-180;
                                  DB 4;
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       Indels 131;
                                  Length 438;
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GENERAL INFORMATION:

APPLICANT: Lederman, Seth

APPLICANT: van Eyndhoven, Winfried

TITLE OF INVENTION: TRAP-3 Deletion Isoforms And Uses Thereof

FILE REFERENCE: 0575-58732

CURRENT APPLICATION NUMBER: US/09/268,544B

CURRENT FILING DATE: 1999-03-11

NUMBER OF SEQ ID NOS: 43

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 38

LENGTH: 347

TYPE: PRT

ORGANISM: Human
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                                                                                                                                                                   Matches 331; Conservative
                                                                                                                                                                                     Query Match
Best Local
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120 GRGCAEQLTLGHLLVHLKNECQFEELPCLRADCKEKVLRKDLRDHVEKACKYREATCSHC 179
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                                                                                                                                                                                  Local Similarity
                                                                                            1 MESSKKMDSPGALQTNPPLKLHTDRSAGTPVFVPEQGGYKEKFVKTVEDKYKCEKCHLVL 60
                                                                                                                1 MESSKKMDAAGTLQPNPPLKLQPDRGAGS-VLVPEQGGYKEKFVKTVEDKYKCEKCRLVL 59
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                                                                                                                                                                           53.6%; Score 1604; DB 4; Length 347; 58.3%; Pred. No. 1.1e-133;
                                                                                                                                                            4; Mismatches 11; Indels 222;
                                                                                                                                                          Gaps
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LENGTH: 558
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: TRAF FAMILY MOLECULES, POLYNUCLEOTIDES ENCODING THEM, TITLE OF INVENTION: AND ANTIBODIES AGAINST THEM FILE REFERENCE: 007898-0255515
CURRENT APPLICATION NUMBER: U5/09/138,277C
CURRENT FILING DATE: 1998-08-18
PRIOR APPLICATION NUMBER: PCT/JP97/00512
PRIOR FILING DATE: 1997-02-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: NAKATA, MOTOMI
APPLICANT: NAKANO, HIROY/
APPLICANT: YAGITA, HIDEO
APPLICANT: OKUMURA, KO
                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: JP 34674/1996 PRIOR FILING DATE: 1996-02-22 NUMBER OF SEQ ID NOS: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: PatentIn Ver.
                                                                                                                                                                                                                                                                                                                                                                                TYPE: PRT
152 DVKEHLSAYCRFREEKCLYCKRDIVVTNLQDHEENSCPAYPVSCPNRC-VQTIPRARVNE 210
                       160 DLRDHVEKACKYREATCSHCKSQVPMIKLQKHEDTDCPCVVVSCPHKCSVQTLLRSELSA 219 | :::|: |::| | ::: |:::| |::::| |::::| |::::|
                                                                                                                    100 FKDNCCKREILALQVYCRNEGRGCAEQLTLGHLLVHLKNECQFEELPCLRADCKEKVLRK 159
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                                                                             94 FKDNCCKREVLNLHVYCKN-APGCNARIILGRFQDHLQH-CSFQAVPCPNESCREAMLRK 151
                                                                                                                                                             34 QEVEQLEERYKCAFCHSVLHNPHQTGCGHRFCQQCIRSLRELNSVPICPVDKEVIKPQEV 93
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YAGITA, HIDEO
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                                                                                                                                                                                                                                                   Conservative 106;
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                                                                                                                                                                                                                                                                  41.5%; Score 1243; DB 4; Length 558; 43.2%; Pred. No. 1.7e-101;
                                                                                                                                                                                                                                                 Mismatches 158;
                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                            48; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ----- 99
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                                                                                                                                                                                                                                                                                                                                                                                                        ; LENGTH: 557
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-138-277C-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: 007988-025515

CURRENT APPLICATION NUMBER: US/99/138,277C

CURRENT FILING DATE: 1980-08-18

PRIOR APPLICATION NUMBER: PCT/JP97/00512

PRIOR FILING DATE: 1997-02-24

PRIOR PILING DATE: 1996-02-22

PRIOR FILING DATE: 1996-02-22

NUMBER OF SEQ ID NOS: 16
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                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 232; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Ver. 2.1 SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: NAKATA, MOTOMI
APPLICANT: NAKANO, HIROYASU
APPLICANT: YAGITA, HIDEO
APPLICANT: OKUMURA, KO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: TRAF FAMILY MOLECULES, POLYNUCLEOTIDES ENCODING TITLE OF INVENTION: AND ANTIBODIES AGAINST THEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         211 HLTVCPEAEQDCPFKHYGCTVKGKRGNLLEHERAALQDHMLLVLEKNYQLEQRISDLYQS 270
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          331 QIVNQQPSRL------QRLVL 368
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   558 IVDTSDLPD 566
                                                                                160 DLRDHVEKACKYREATCSHCKSQVPMIKLQKHEDTDCPCVVVSCPHKCSVQTLLRSELSA 219
                                                                                                                                                                          100 FKDNCCKREILALQVYCRNEGRGCAEQLTLGHLLVHLKNECQFEELPCLRADCKEKVLRK 159
                                                                                                                                                                                                                                                   41 KFVKTVEDKYKCEKCRLVLCNPKQTECGHRFCESCMAALLS-SSSPKCTACQESIIKDKV 99
                                                                                                                                                                                                                         34 QFVERLEERYKCAFCHSVLHNPHQTGCGHRFCQHCILSLRELNTVPICPVDKEVIKSQEV 93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LEQKESKIQQLAETVKKFEKELKQFTQMFGRNGTFLSNVQALTSHTDKSAWLEAQVRHLL 330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SVEKNKSIQSLHNQICSFEIEIEROKEMLRNN------ESKILHLQ 319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HLSECVNAPSTCSFKRYGCVFQGTNQQIKAHEASSAVQHVNLLKEWSNSLEKKVSLLQNE 279
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        YSQPFYTGYFGYKMCARVYLNGDGMGKGTHLSLFFVIMRGEYDALLPWPFKQKVTLMLMD 499
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RVIDSQAEKLKELDKEIRPFRONWEEADSMKSSVESLONRVTELESVDKSAGQAARNTGL 379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AVDLTDLED 557
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     QGSSRRHLGDAFKPDPNSSSFKKPTGEMNIASGCPVFVAQTVLEN--GTYIKDDTIFIKV 557
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FSQPFYTSRCGYRLCARAYLNGDGSGKGTHLSLYFVVMRGEFDSLLQWPFRQRVTLMLLD 488
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LESQLSRHDQTLSVHDIRLADMDLRFQVLETASYNGVLIWKIRDYKRRKQEAVMGKTLSL 439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LEGETSKHDAHINIHKAQLNKNEERFKQLEGACYSGKLIWKVTDYRVKKREAVEGHTVSV 428
                                        DLKEHLSASCOFRKEKCLYCKKDVVVINLQNHEENLCPEYPVFCPNNCA-KIILKTEVDE 210
                                                                                                                                      HLSECVNAPSTCSFKRYGCVFQGTNQQIKAHEASSAVQHVNLLKEWSNSLEKKVSLLQNE
                                                                                                                                                                                                                                                                                                                     Conservative 112; Mismatches
                                                                                                                                                                                                                                                                                                                                          39.7%; Score 1189.5; DB 4; Length 557; 43.4%; Pred. No. 9.3e-97;
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US-09-181-958-2
RESULT 8
US-08-331-394-4
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LENGTH: 228
TYPE: PRT
ORGANISM: human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 2, Application US/09181958 Patent No. 6143507
                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: High Throughput Compatible Assays for Receptor-TRAF TITLE OF INVENTION: Interactions FILE REFERENCE: 9_142_Nucl.Seq CURRENT APPLICATION NUMBER: US/09/181,958 CURRENT FILING DATE: 198-10-29 NUMBER OF SEQ ID NOS: 2 SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Kehry, Marilyn R
APPLICANT: Pullen, Steven S
APPLICANT: Crute, James J
APPLICANT: Crute, James J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 514 DPNSSSFKKPTGEMNIASGCPVFVAQTVLEN--GTYIKDDTIFIKVIVDTSDLPD 566
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            383 HKAQLSKNEERFKLLEGTCYNGKLIWKVTDYKMKKREAVDGHTVSIFSQSFYTSRCGYRL 442
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     394 HDIRLADMDLRPQVLETASYNGVLIWKIRDYKRRKQEAVMGKTLSLYSQPFYTGYFGYKM 453
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      330 LQMVNQQQNKFDLRPLMEAVDTVKQKTTLLENND-----QRLAVLEEETNKHDTHINI 382
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        340 -----RONWEEADSMKSSVESLONRVTELESVDKSAGQAARNTGLLESQLSRHDQTLSV 393
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             271 LEQKESKIQQLAETIKKLEKEFKQFAQLFGKNGSFLPNIQ-VFASHIDKSAWLEAQVHQL 329
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               280 SVEKNKSIQSLHNQICSFEIEIERQKEMLRNNESKILHLQRVIDSQAEKLKELDKEIRPF 339
                                                                                                                                                                                                                                                                                                                     353 VESLQNRVTELESVDKSAGQAARNTGLLESQLSRHDQTLSVHDIRLADMDLRFQVLETAS 412
                                                                                                                                                    121
                                                                                                                                                                                      473
                                                                                                                                                                                                                      61 YNGVLIWKIRDYKRRKQEAVMGKTLSLYSOPFYTGYFGYKMCARVYLNGDGMGKGTHLSL 120
                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
nes 213; Conserv
                                                                                                                                                                                                                                                                                             1 VESLQNRVTELESVDKSAGQVARNTGLLESQLSRHDQMLSVHDIRLADMDLRFQVLETAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DPNSSSFKRPDGEMNIASGCPRFVAHSVLENAKNAYIKDDTLFLKVAVDLTDLED 556
                                                                            CPVFVAQTVLENGTYIKDDTIFIKVIVDTSDLPDP 567
                                                                                                                                                FFVIMRGEYDALLPWPFKQKVTLMLMDQGSSRRHLGDAFKPDPNSSSFKKPTGEMNIASG
                                                                                                                                                                     FFVIMRGEYDALLPWPFKQKYTLMLMDQGSSRRHLGDAFKPDPNSSSFKKPTGEMNIASG
                                                                                                                                                                                                                                         YNGVLIWKIRDYKRRKQEAVMGKTLSLYSQPFYTGYFGYKMCARVYLNGDGMGKGTHLSL 472
                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                        37.0%;
99.1%;
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Pred. No. 4e-90;
                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                          2;
                                                                                                                                                                                                                                                                                                                                                                                                           Length 228;
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                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                           180
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Sequence 4, Application US/08331394 Patent No. 5670319

GENERAL INFORMATION:
APPLICANT: Goedde

APPLICANT:

Rothe, Mike

Goeddel, David V.

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INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 501 amino
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
344 ELEVSTYDGVFIWKISDFTRKRQEAVAGRTPAIFSPAFYTSRYGYKMCLRVYLNGDGTGR 403
                                      407 VLETASYNGVLIWKIRDYKRRKQEAVMGKTLSLYSQPFYTGYFGYKMCARVYLNGDGMGK 466
                                                                                                                                        347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-3216
TELEFAX: 415/952-9881
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/250858
FILING DATE: 27-MAY-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-SOTTWARE: patin (Genentech) CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 66
CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STATE: California
                                                                                                                                                                                                                                                   222 GCSEMVETENLQDHELQRLREHLALL--LSSFLEAQASPGTLNQVGPELLQRCQILEQK- 278
                                                                                                                                                                                                                                                                                              237 GCVFQGTNQQIKAHEASSAVQHVNLLKEWSNSLEKKVS------LLQNESVEKNKS 286
                                                                                                                                                                                                                                                                                                                                        164 QHCRAPCSHVDLEVHYEV-CPKFPLTC-DGCGKKKIPRETFQDHVRACSKCRVLCRFHTV 221
                                                                                                                                                                                                                                                                                                                                                                            177 SHCKSQVPMIKLQKHEDTDCPCVVVSCPHKCSVQTLLRSELSAHLSECVNAPSTCSFKRY 236
                                                                                                                                                                                                                                                                                                                                                                                                                         108 PND--GCTWKGTLKEYESCHEGLCPFLLTEC--PACKGLVRLSEKEHHTEQECPKRSLSC 163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            48 QCGHRYCSFCLTSILSSGPQNCAACVYEGLYEEGISILESSSAFPDNAARREVESLPAVC 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 MAAASVTSPGSLELLQP------GFSKTLLGTRLEAKYLCSACKNILRRPPQA 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7 MDAAGTLQPNPPLKLQPDRGAGSVLVPEQGGYKEKFVKT-VEDKYKCEKCRLVLCNPKQT 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Dreger, Ginger R. REGISTRATION NUMBER: 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: 5.25 inch, 360 Kb floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                           DSMKSSVESLQNRVTELESVDKSAGQAARNTGLLESQLSRHDQTLSVHDIRLADMDLRFQ 406
                                                                                                                                                                                                             IQSLHNQICSFEIEIERQKEMLRNNESKILHLQRVIDSQAEKLKELDKEIRPFRQNWEEA 346
                                                                                     ----KIEALSNKVQQLE------RSIGLKDLAMADLEQKVS
                                                                                                                                                                            IATFENIVCVLNREVER--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RNEGRGCAEQLTLGHLLVHLKNECQFEELPCLRADCKEKVLRKDLRDHVEKACKYREATC 176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ECGHRFCESCMAALLSSSSPKCTAC-----QESII----KDKVFKDNCCKREILALQVYC 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        194;
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US-08-250-858-4
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                                                                                                                                                                                                                                                                                                                                                                          Matches 194;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 4, Application US/08250858
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: 89
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-3216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Ginger R.
REGISTRATION NUMBER: 33,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
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PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Goeddel, David V.
APPLICANT: Rothe, Mike
TITLE OF INVENTION: Tumor Necrosis Factor Receptor-Associated Factors
NUMBER OF SEQUENCES: 62
                                        164 QHCRAPCSHVDLEVHYEV-CPKFPLTC-DGCGKKKIPRETFQDHVRACSKCRVLCRFHTV 221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
237 GCVFQGTNQQIKAHEASSAVQHVNILKEWSNSLEKKVS------LLQNESVEKNKS 286
                                                                                                                      108 PND--GCTWKGTLKEYESCHEGLCPFLLTEC--PACKGLVRLSEKEHHTEQECPKRSLSC 163
                                                                                                                                                            117 RNEGRGCAEQLTLGHLLVHLKNECQFEELPCLRADCKEKVLRKDLRDHVEKACKYREATC 176
                                                                                                                                                                                                               48
                                                                                                                                                                                                                                           66 ECGHRFCESCMAALLSSSSPKCTAC-----QESII---KDKVFKDNCCKREILALQVYC 116
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 417, 7168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/0 FILING DATE: 27-May-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                463 MNIASGCPLFCPVSKMEAKNSYVRDDAIFIKAIVDLTGL 501
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: patin (Genentech)
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CITY: South San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                       7 MDAAGTLQPNPPLKLQPDRGAGSVLVPEQGGYKEKFVKT-VEDKYKCEKCRLVLCNPKQT 65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH:
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                                                                              SHCKSQVPMIKLQKHEDTDCPCVVVSCPHKCSVQTLLRSELSAHLSECVNAPSTCSFKRY 236
                                                                                                                                                                                                    QCGHRYCSFCLTSILSSGPQNCAACVYEGLYEEGISILESSSAFPDNAARREVESLPAVC 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          amino acid
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                                                                                                                                                                                                                                                                                                                                                                   28.0%; Score 837.5; 33.5%; Pred. No. 1e-
tive 88; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                 core 837.5; DB 1; red. No. 1e-65; Mismatches 198;
                                                                                                                                                                                                                                                                                                                                                                                                       Length 501;
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Matches
            Query Match
Best Local Similarity
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                                                                                                                                                      TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: 5.25 inch, 360 Kb floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Tumor Necrosis Factor Receptor-Associated Factors NUMBER OF SEQUENCES: 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Goeddel, Dav
APPLICANT: Rothe. Mike
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                                                                                                                                                                                                                                                                                            FILING DATE: 28-OCT-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     404
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                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
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OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE:
                                                                                                                                                                                                                                                             NAME: Dreger, Ginger R. REGISTRATION NUMBER: 33
                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 08/21 FILING DATE: 27-MAY-1994
                                                                                                    TYPE:
                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
                                                                                    TOPOLOGY:
                                                                                                                     LENGTH:
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                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER:
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South San Francisco
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Conservative
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                28.0%; Score 837.5; 33.5%; Pred. No. 1e-
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                837.5; DB 1;
No. 1e-65;
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                                  Length 501;
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US-08-744-139-4
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                                                                                                                                   SOFTWARE: WinPatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION UNMEER: US/08/744,
FILING DATE: 31-Oct-1996
                  APPLICATION NUMBER: 08/25
FILING DATE: 05/27/1994
ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Ginger R.
                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
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NUMBER OF SEQUENCES:
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                                                                                              CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    279
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                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Genentech, Inc.
STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7 MDAAGTLQPNPPLKLQPDRGAGSVLVPEQGGYKEKFVKT-VEDKYKCEKCRLVLCNPKQT
                                                                                                                                                                                                              COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                                                                                                                                                                                        MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           QHCRAPCSHVDLEVHYEV-CPKFPLTC-DGCGKKKIPRETFQDHVRACSKCRVLCRFHTV 221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         VLETASYNGVLIWKIRDYKRRKQEAVMGKTLSLYSQPFYTGYFGYKMCARVYLNGDGMGK 466
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GCSEMVETENLODHELORLREHLALL--LSSFLEAQASPGTLNQVGPELLORCQILEQK- 278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RNEGRGCAEQLTLGHLLVHLKNECQFEELPCLRADCKEKVLRKDLRDHVEKACKYREATC 176
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UMBER: 33,055
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                                                                                                                                                                                                        Sequence 4, Application PC/TUS9506639 GENERAL INFORMATION:
APPLICANT: Genentech, Inc.
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COMPUTER READABLE FORM: MEDIUM TYPE: 3.5 inch,
                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STATE: California
                                                                                                                                                               APPLICANT: Genentech, Inc.
TITLE OF INVENTION: Tumor Necrosis Factor Receptor-Associated
NUMBER OF SEQUENCES: 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 501 amino acids
TYPE: Amino Acid
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                                       ZIP:
                                                       COUNTRY:
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TELEFAX: 910/371-7168
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER:
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 1.44 Mb
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TELEPHONE: 415/225-3216
TELEFAX: 415/952-9881
TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 28-OCT-1994 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
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LENGTH: 501 amino acids
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                                    527 MNIASGCPVFVAQTVLE-NGTYIKDDTIFIKVIVDTSDL 564
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TYPE: amino acid
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
CLASSIFICATION:
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MNIASGCPLFCPVSKMEAKNSYVRDDAIFIKAIVDLTGL 501
                                                                                                     GTHLSLFFVIMRGEYDALLPWPFKQKVTLMLMDQGSSRRHLGDAFKPDPNSSSFKKPTGE 526
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                                                                               GTHLSLFFVVMKGPNDALLQWPFNQKVTLMLLDH-NNREHVIDAFRPDVTSSSFQRPVSD
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US-08-331-394-2
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APPLICANT: Goedde
APPLICANT: Rothe,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 409 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 89
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-3216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOTTWARE: patin (Genentech) CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Tumor Necrosis Factor TITLE OF INVENTION: Receptor-Associated Factors NUMBER OF SEQUENCES: 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 08/25
FILING DATE: 27-MAY-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: 5.25 inch,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     137 KNECQFEELPCLRADCKEKVLRKDLRDHVEKACKYREATCSHCKSQVPMIKLQKHEDTDC 196
                                                                                                                                       201 VE-----ASHLALAASIHQSQLDREHLLSLEQRVVELQQTLAQKDQVLGKLEHSLRLM 253
                                                                                                                                                                                     361 TELESVDKSAGQAARNTGLLESQLSRH-------DQTLSVHDIRLADMDLRFQVL 408
                                                                                                                                                                                                                                                                            302 ERQKEMLRNNESKILHLQRVIDSQAEK-LKELDKEIRPFRQNWEEADSMKSSVESLQNRV 360
                                                                                                                                                                                                                                158 YRAPCCESQEELALQHLVK-----EKLLAQLEEKLRVF------ANIVAVLNKE 200
                                                                                                                                                                                                                                                                                                                                                                     257 QHVNL----LKEWSNS------LEKKVSLLQNESVEKNKSIQSLHNQICSFEIEI 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 460 Point San Bru
CITY: South San Francisco
STATE: California
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                                                                                                                                                                                                                                                                                                                                                                                                                     49 PKCRADNLHPVSPGSPLTQE-KVH-SDVAEAEIMCPFAGVGCSFKGSPQSMQEHEATSQS 106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 415/952-9881
TELEX: 910/371-7168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Dreger, Ginger R. REGISTRATION NUMBER: 33,055
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            9
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es 162; Conserv
HLSLFFVIMRGEYDALLPWPFKQKVTLMLMDQGSSRRHLGDAFKPDPNSSSFKKPTGEMN 528
                                                EEASFDGTFLWKITNVTKRCHESVCGRTVSLFSPAFYTAKYGYKLCLRLYLNGDGSGKKT 313
                                                                                         ETASYNGVLIWKIRDYKRRKQEAVMGKTLSLYSQPFYTGYFGYKMCARVYLNGDGMGKGT 468
                                                                                                                                                                                                                                                                                                                         SHLYLLLAVLKEWKSSPGSNLGSAPMALERNLSELQ------LQAAVEATGDLEVDC 157
                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCVVVSCPHKCSVQTLLRSELSAHLSECVNAPSTCSFKRYGCVFQGTNQQIKAHEASSAV 256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ENEFQF---GCPPAPCQD------PSEPRVLCCTACLSE----NLRDDEDRIC 48
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21.4%; Score 640; DB 1; Length 409; 35.5%; Pred. No. 2.1e-48; tive 72; Mismatches 136; Indels
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RESULT 14
US-08-250-858-2
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                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM: PC-DOS/MS
SOFTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/250,858 FILING DATE: 27-May-1994 CLASSIFICATION: 435 PRIOR APPLICATION DATA:
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MEDIUM TYPE: 5.25 inch,
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                 361 TELESVDKSAGQAARNTGLLESQLSRH-----
                                                   158 YRAPCCESQEELALQHLVK-----EKLLAQLEEKLRVF------ANIVAVLNKE 200
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                                                                                                                                                                                                                                                                                                        137 KNECQFEELPCLRADCKEKVLRKDLRDHVEKACKYREATCSHCKSQVPMIKLQKHEDTDC 196
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Dreger, Ginger REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE
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                                                                                  ERQKEMLRNNESKILHLQRVIDSQAEK-LKELDKEIRPFRQNWEEADSMKSSVESLQNRV 360
                                                                                                                          SHLYLLLAVLKEWKSSPGSNLGSAPMALERNLSELQ------LQAAVEATGDLEVDC 157
                                                                                                                                                          QHVNL----LKEWSNS------LEKKVSLLQNESVEKNKSIQSLHNQICSFEIEI 301
                                                                                                                                                                                                PKCRADNLHPVSPGSPLTQE-KVH-SDVAEAEIMCPFAGVGCSFKGSPQSMQEHEATSQS 106
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910/371-7168
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SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                              72;
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-----DQTLSVHDIRLADMDLRFQVL 408
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US-08-446-915-2
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/250
FILING DATE: 27-MAY-1994
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                            TELEX: 910/371-7168
INFORMATION FOR SEQ ID NO: 2:
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                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 409 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                              NAME: Dreger, Ginger R. 055.
REGISTRATION NUMBER: 33.055.
REFERENCE/DOCKET NUMBER: 89.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415/225-3216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: 5.25 inch, 360 kb floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: patin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/446,915
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 08/331394
FILING DATE: 28-OCT-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Genentech, Inc.
STREET: 460 Point San Bruno Blvd
CITY: South San Francisco
STATE: California
                                 197 PCVVVSCPHKCSVQTLLRSELSAHLSECVNAPSTCSFKRYGCVFQGTNQQIKAHEASSAV 256
                                                                                                          137 KNECOFEELPCLRADCKEKVLRKDLRDHVEKACKYREATCSHCKSQVPMIKLQKHEDTDC 196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Goeddel, David v.
APPLICANT: Rothe. Mike
TITLE OF INVENTION: Tumor Necrosis Factor Receptor-Associated Factors
NUMBER OF SEQUENCES: 66
49 PKCRADNLHPVSPGSPLTQE-KVH-SDVAEAEIMCPFAGVGCSFKGSPQSMQEHEATSQS 106
                                                                                                                                                         Local Similarity 35.5 les 162; Conservative
                                                                                                                                                                                                                                                                                                                                                TELEPHONE: 415/252-9881
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             469 HLSLFFVIMRGEYDALLPWPFKOKVTLMLMDQGSSRRHLGDAFKDDPNSSSFKKPTGEMN 528
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       USA
                                                                                                                                                                                                                                                        linear
                                                                                                                                                 21.4%; Score 640; DB 1; Length 409; 35.5%; Pred. No. 2.1e-48; tive 72; Mismatches 136; Indels
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                                                                                                                                                 86; Gaps
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Qy 529 IASGCPVFVAQTVLEN	Qy 469 HLSLFFVIMRGEYDAI 	Qy 409 ETASYNGVLIWKIRDY       ::  :    :   Db 254 EEASFDGTFLWKITNV	Qy 361 TELESVDKSAGQAARN	Qy 302 ERQKEMLRNNESKILF 	Qy 257 QHVNLLKEWSNS-  :                Db 107 SHLYLLLAVLKEWKSSF
529 IASGCPVEVAQTVLENGTYIKDDTIFIKVIVDTS 562 	469 HLSLFFVIMRGEYDALLPWPFKQKVTLMLMDQGSSRRHLGDAFKPDPNSSSFKKPTGEMN 528	409 ETASYNGVLIWKIRDYKRRKQEAVMGKTLSLYSQPFYTGYFGYKMCARVYLNGDGMGKGT 468 	361 TELESVDKSAGQAARNTGLLESQLSRH	302 ERQKEMLRNNESKILHLQRVIDSQAEK-LKELDKEIRPFRQNWEEADSMKSSVESLQNRV 360	257 QHVNLLKEWSNSLEKKVSLLQNESVEKNKSIQSLHNQICSFEIEI 301  :
	528 372	468 313	408 253	360 200	301 157

Search completed: December 19, 2002, 15:02:21 Job time : 18 secs